

SEQUENCE LISTING

10/510677

SEQ ID NO.: 1; AAC2-1 nucleotide sequence

5 AC  
 TCGCCACTCCTCCGACGTGCTGGGCAACCTCAACGAGCTGCGCCTGCGCGGGATCCTCACTGACGTCACGCTGCT  
 GG  
 TTGGCGGGCAACCCCTCAGAGCACACAAGGCAGTTCTCATCGCCTGCAGTGGCTTCTTCTATTCAATTTTCCGGG  
 GC  
 10 CGTGCGGGAGTCGGGGTGGACGTGCTCTCTCTGCCCCGGGGTCCCGAAGCGAGAGGCTTCGCCCCCTCTATTGGAC  
 TT  
 CATGTACACTTCGCGCCTGCGCCTCTCTCCAGCCACTGCACCAGCAGTCCTAGCGGCCGCCACCTATTTGCAGAT  
 GG  
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 15 AA  
 GCAGAACCCCCAACACCCCCAACGGCCCCCTCCACCAGGTAGTCCAGGCGCTCCGAAGGACACCCAGACCCACCT  
 AC  
 TGAATCTCGAAGCTGCAGTCAAGGCCCCCAGTCCAGCCAGCCCTGACCCCAAGGCCTGCAACTGGAAAAAGTA  
 CA  
 20 AGTACATCGTGCTAAACTCTCAGGCCTCCCAAGCAGGGAGCCTGGTTCGGGGAGAGAAGTTCTGGTCAACCTTGCC  
 CC  
 CAAGCCAGGCTCCCCAGTGGAGACGAGGCCTCCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGTGAAGAAGGA  
 CC  
 CATTCCTGGTCCCCAGAGCAGGCTCTCTCCAAGTCTGCCACTGTGCAGTTCAAATGTGGGGCTCCAGCCAGTAC  
 25 CC  
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 AT  
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 GA  
 30 CAAACCTATAAGTGTGAGCTGTGCCGGTCTTCGTTCCGCTACAAGGGCAACCTTGCCAGTCACCGTACAGTGCA  
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 GC  
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 35 CA  
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 CA  
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 GT  
 40 CAACTGCGGCTGCATCTGCGCCAGAAACACGGAGCTGCTACCAACACCAAAGTGCACTACCACATTCTCGGGGGG  
 CC  
 CTAGCTGAGCGCAGGCCAGGCCCACTTGCTTCCTGCGGGTGGGAAAGCTGCAGGCCCAGGCCTTGCTTCCCTA  
 TC  
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 45 GC  
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 GC  
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 CT  
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 AA

CCCTCTCTGGTATTCTGGATGTTGTAGGTTCTCTAGCAGTCTAGAAATGGATACAGACATTTCTCTGTTCTTCAA  
GG  
GTGATAGGAACCATTATGTTGAGCCCAAAATGGAAGTAATAATAAATGCCTCCTGGAGGCTGTGGGTGTGGGGGA  
TT  
5 CTGTATCTGGATTCCGTATCACTCCAACCTGGAGGCTGTGGGTGTGGGGGATTCTGTATCTGGATTCCGTATCACT  
CC  
AAGTGGAGGCTGGCAGGTTTTTCTGCAAGATGGTCCAGAATCTAAAATGTCCCATTAATCTGGTCACTTGGGTTT  
GG  
CTCTGCTGTATCCATCTATAGTGGTAGAGACCCACCAGGGCTCAAGTGGAGTCCATCATCCTCCCACGGGGGCCT  
10 GT  
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CT  
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GT  
15 TAACCCATCCTTTACTACAGAGGCATATGGGTTTGAATGTTACCTGGGGTTCTCTCTATTGAGTTGAGCCCCCTC  
TT  
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CT  
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20 AT  
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AT  
25 TAAGAGGTTGGTTGAGGGGTGCAGTTTCTGGTGTAGGCCAGGTAGGTAGAAAGTGAGGAACAGGGTTGCCTCTTG  
GC  
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TT  
ATCTGATTATGGGACGAGGGTAGAAAGTAAGAAGCACTTTTGAATTTGTGGGGTAGAACTTCAACAATAAGTCAG  
30 TT  
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GC  
AGAATTAAAGAAGGAAGAAGGAAGGCGGAGGAGTCTATAAGAAGGAATCATGATTTCTATTTAGCAGATTGGAT  
GG  
35 GCAGGTGGAGAATGCCTGGGGGTAGAAATGTTAGATCTTGCAACATCAGATCCTTGAATAAAGAAGCCTCTCTG  
CG  
CAAAAAAAAAAAAAAAAAAAAAA

SEQ ID NO.: 2; AAC2-2 Open reading frame

ATGGGTTCCCCCGCGCCCGGAGGGAGCGCTGGGCTACGTCCGAGTTCACTCGCCACTCCTCCGACGTGCTGGGCAACCT  
CAACGAGCTGCGCCTGCGCGGGATCCTCACTGACGTACGCTGCTGGTTGGCGGGCAACCCCTCAGAGCACACAAGGCAGTTC  
5 TCATCGCCTGCAGTGGCTTCTTCTATTCAATTTTCCGGGGCCGTGCGGGAGTCGGGGTGGACGTGCTCTCTCTGCCCCGGGGT  
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CCTAGCGGCGCCACCTATTTGCAGATGGAGACGTGGTCCAGGCATGCCACCGCTTCATCCAGGCCAGCTATGAACCTCTGG  
GCATCTCCCTGCGCCCCCTGGAAGCAGAACCCCCAACACCCCCAACGGCCCCCTCCACCAGGTAGTCCAGGCGCTCCGAAGGA  
10 CACCAGACCCACCTACTGAATCTCGAAGCTGCAGTCAAGGCCCCCAGTCCAGCCAGCCCTGACCCCAAGGCCTGCAACTG  
GAAAAAGTACAAGTACATCGTGCTAACTCTCAGGCCTCCCAGCAGGGAGCCCTGGTGGGGGAGAGAAGTTCTGGTCAACCTT  
GCCCCAAGCCAGGCTCCCCAGTGGAGACGAGGCCTCCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGTGAAGAAGGACCCATT  
CCTGGTCCCCAGAGCAGGCTCTCTCCAATGCTGCCACTGTGCAGTTCAAATGTGGGGCTCCAGCCAGTACCCCTACCTCCT  
CACATCCCAGGCTCAAGACACCTCTGGATCACCTCTGAACGGGCTCGTCCACTACCGGGAAGTGAATTTTTCAGCTGCCAGA  
15 ACTGTGAGGCTGTGGCAGGGTGTCTATCGGGGCTGGACTCCTTGGTTCTGGGGACGAAGACAAACCCTATAAGTGTGCACTG  
TGCCGGTCTTCGTTCCGCTACAAGGGCAACCTTGCCAGTCATCGTACAGTGCACACAGGGGAAAAGCCTTACCACTGCTCAAT  
CTGCGGAGCCCGTTTAAACCGGCCAGCAAACTGAAAACGCACAGCCGCATCCATTGCGGAGAGAAGCCGTATAAGTGTGAGA  
CGTGCGGCTCGCGCTTTGTACAGGTGGCACATCTGCGGGCGCAGGTGCTGATCCACACCGGGGAGAAGCCCTACCTTGGCCCT  
20 ACCTGCGGAACCCGCTTCCGCCACCTGCAGACCTCAAGAGCCAGTTTCGATCCACACCGGAGAGAAGCCTTACCACTGCGA  
CCCCTGTGGCCTGCATTTCGGGCACAAGAGTCAACTGCGGCTGCATCTGCGCCAGAAACAGGAGCTGCTACCAACACCAAAG  
TGCACTACCACATTCTCGGGGGGCCCTAG

SEQ ID NO.: 3; 7524

ATACCCGGAACCTCCCTAAGCCTTCTATTAGCTCCAATAATAGTAAGCCTGTGCAAGACAAAGATG

SEQ ID NO.: 4; 7526

GCCTGTGTCCCCTAGACTCCAACCTCAGCAACGGAAATAGAACTCTGACCCTGTTAACGTGACCAGGAAC

SEQ ID NO.: 5; 7528

ACGTGCTTTACGGACCCGATGCTCCTACAATCAGCCCTCTAAACACAAGCTATAGATCAGGGGAAAATCT

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SEQ ID NO.: 6; 7533

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SEQ ID NO.: 7; 7535

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SEQ ID NO.: 8; 7537

GATCCACTATTGTTTCACGGTAATATTGGGAATGAACAGTTCCTGGGTGGACTGTTGGAAAGTG

SEQ ID NO.: 9; 7567

GACACAGCAAGCTACAAATGCGAAACCCAAAATCCAGTCAGCGCCAGGAGGTCTGATTCACTGATTCTCA

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SEQ ID NO.: 10; 7568

TGAATCAGACCTCCTGGCGCTGACTGGATTTTGGGTTTCGCATTTGTAGCTTGCTGTGTCGTTCCCTGGTC

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SEQ ID NO.: 11; 7576

GATCCTACACGTGCCAAGCTCACAATAGCGACACCGGACTCAACCGCACAAACCGTGACGACGATTACCGTGTATG  
CCGA

SEQ ID NO.: 12; 7587

CATCCTCAACTGGGTTAGAATTGTTACTAGTTATGAATGGTTTTGGTGGCTCGGCATACACGGTAATCGT

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SEQ ID NO.: 13; 7677

TTCTAACCAGTTGAGGATGAGGACGCAGTTGCATTAACTTGTGAGCCAGAGATTCAAAATACCACTTATTTATG  
55 GTGGG

SEQ ID NO.: 14; 7678

GTCTAATGATAACCGCACATTGACACTCCTGTCCGTTACTCGCAATGATGTAGGACCTTATGAGTGTGGCATTCA  
GAATG

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SEQ ID NO.: 15; 7679

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TTGCC

5 SEQ ID NO.: 16; 7680

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TTGCC

10 SEQ ID NO.: 17; 7681

TAAACAATAACTGTTTCCGCGGAGCTGCCCAAGCCCTCCATCTCCAGCAACAACTCCAAACCCGTGGAGGACAA  
GGATG

15 SEQ ID NO.: 18; 7682

ATGTGCGGTTATCATTAGACAACTGCAAGCGTGGGCTAACCGGCAAACCTTTGGTTATTGACCCACCATAAATAAG  
TGGTA

20 SEQ ID NO.: 19; 7683

GGTCGTCTGGGCCATACAAAACATTAAGGATAACAGGGTCGGAGTGATCAACGGATAATTCACTTCTGAATGCCAC  
ACTCA

25 SEQ ID NO.: 20; 7684

GCTGCTGAATGTTTCCATCAATCAGCCAGGAGTACTGTGCAGGGGGGTGGATGCTGCATGGCAAGAAAGGCTCA  
AGTTC

30 SEQ ID NO.: 21; 7685

CGGAAACAGTTATTGTTTTAACTGTAGTCCTGCTGTGACCACTGGCTGAGTTATTGGCCTGGCAAGTATAGAGTC  
CGCTG

35 SEQ ID NO.: 22; 7686

CCTCAGGTTTCACAGGTGAAGGCCACAGCATCCTTGTCTCCACGGGT

SEQ ID NO.: 23; CEA-CAP6D

ATGGAGTCTC CCTCGGCCCC TCCCCACAGA TGGTGCATCC CCTGGCAGAG GCTCCTGCTC  
ACAGCCTCAC TTCTAACCTT CTGGAACCCG CCCACCACTG CCAAGCTCAC TATTGAATCC  
35 ACGCCGTTCA ATGTCGCAGA GGGGAAGGAG GTGCTTCTAC TTGTCCACAA TCTGCCCCAG  
CATCTTTTTG GCTACAGCTG GTACAAAGGT GAAAGAGTGG ATGGCAACCG TCAAATTATA  
GGATATGTAA TAGGAACCTA ACAAGCTACC CCAGGGCCCC CATAAGTGG TCGAGAGATA  
ATATACCCCA ATGCATCCCT GCTGATCCAG AACATCATCC AGAATGACAC AGGATTCTAC  
ACCCACACAG TCATAAAGTC AGATCTTGTG AATGAAGAAG CAACTGGCCA GTTCCGGGTA  
40 TACCCGAGC TGCCCAAGCC CTCCATCTCC AGCAACAAC CCAAACCCGT GGAGGACAAG  
GATGCTGTGG CTTTACCTG TGAACCTGAG ACTCAGGACG CAACCTACCT GTGGTGGGTA  
AACAATCAGA GCCTCCCGGT CAGTCCAGG CTGCAGCTGT CCAATGGCAA CAGGACCCCTC  
ACTCTATTCA ATGTCACAAG AAATGACACA GCAAGCTACA AATGTGAAAC CCAGAACCCA  
GTGAGTGCCA GGCGCAGTGA TTCAGTCATC CTGAATGTCC TCTATGGCCC GGATGCCCCC  
45 ACCATTTCCC CTCTAAACAC ATCTTACAGA TCAGGGGAAA ATCTGAACCT CTCCTGCCAC  
GCAGCCTCTA ACCACCTGC ACAGTACTCT TGGTTTGTCA ATGGGACTTT CCAGCAATCC  
ACCCAAGAGC TCTTTATCCC CAACATCACT GTGAATAATA GTGGATCCTA TACGTGCCAA  
GCCCATAACT CAGACACTGG CCTCAATAGG ACCACAGTCA CGACGATCAC AGTCTATGAG  
CCACCCAAAC CTTTATCAC CAGCAACAAC TCCAACCCCG TGGAGGATGA GGATGCTGTA  
50 GCCTTAACCT GTGAACCTGA GATTCAGAAC ACAACCTACC TGTGGTGGGT AAATAATCAG  
AGCCTCCCGG TCAGTCCAG GCTGCAGCTG TCCAATGACA ACAGGACCCCT CACTCTACTC  
AGTGTACAAA GGAATGATGT AGGACCCTAT GAGTGTGGAA TCCAGAACGA ATTAAGTGTT  
GACCACAGCG ACCCAGTCAT CCTGAATGTC CTCTATGGCC CAGACGACCC CACCATTTC  
CCCTCATACA CCTATTACCG TCCAGGGGTG AACCTCAGCC TCTCCTGCCA TGCAGCCTCT  
55 AACCACCTG CACAGTATTC TTGGCTGATT GATGGGAACA TCCAGCAACA CACACAAGAG  
CTCTTTATCT CCAACATCAC TGAGAAGAAC AGCGGACTCT ATACCTGCCA GGCCAATAAC  
TCAGCCAGTG GCCACAGCAG GACTACAGTC AAGACAATCA CAGTCTCTGC GGAGCTGCCC  
AAGCCCTCCA TCTCCAGCAA CAACTCCAAA CCCGTGGAGG ACAAGGATGC TGTGGCCTTC  
ACCTGTGAAC CTGAGGCTCA GAACACAACC TACCTGTGGT GGGTAAATGG TCAGAGCCTC  
60 CCAGTCAGTC CCAGGCTGCA GCTGTCCAAT GGCAACAGGA CCCTCACTCT ATTCAATGTC

5 ACAAGAAATG ACGCAAGAGC CTATGTATGT GGAATCCAGA ACTCAGTGAG TGCAAACCGC  
 AGTGACCCAG TCACCCTGGA TGTCTCTAT GGGCCGGACA CCCCATCAT TTCCCCCCA  
 GACTCGTCTT ACCTTTCGGG AGCGGACCTC AACCTCTCCT GCCACTCGGC CTCTAACCCA  
 TCCCCGAGT ATTCTTGGCG TATCAATGGG ATACCGCAGC AACACACACA AGTTCTCTTT  
 ATCGCCAAAA TCACGCCAAA TAATAACGGG ACCTATGCCT GTTTTGTCTC TAACTTGGCT  
 ACTGGCCGCA ATAATTCCAT AGTCAAGAGC ATCACAGTCT CTGCATCTGG AACTTCTCCT  
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SEQ ID NO.: 24; CAP6D-1,2

10 ATGGAGTCTC CCTCGGCCCC TCCCCACAGA TGGTGCATCC CCTGGCAGAG GCTCCTGCTC  
 ACAGCCTCAC TTCTAACCTT CTGGAACCCG CCCACCACTG CCAAGCTCAC TATTGAATCC  
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 CATCTTTTGT GCTACAGCTG GTACAAAGGT GAAAGAGTGG ATGGCAACCG TCAAATTATA  
 GGATATGTAA TAGGAACTCA ACAAGCTACC CCAGGGCCCCG CATACAGTGG TCGAGAGATA  
 15 ATATACCCCA ATGCATCCCT GCTGATCCAG AACATCATCC AGAATGACAC AGGATTCTAC  
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 TACCCGGAAC TCCCTAAGCC TTCTATTAGC TCCAATAATA GTAAGCCTGT CGAAGACAAA  
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 20 ACCCTGTTTA ACGTGACCAG GAACGACACA GCAAGCTACA AATGCGAAAC CAAAATCCA  
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 25 GCTCACAATA GCGACACCGG ACTCAACCGC ACAACCGTGA CGACGATTAC CGTGTATGAG  
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 30 GATCACTCCG ACCCTGTTAT CCTTAATGTT TTGTATGGCC CAGACGACCC AACTATATCT  
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 35 AAGCCCTCCA TCTCCAGCAA CAACTCCAAA CCCGTGGAGG ACAAGGATGC TGTGGCCTTC  
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 40 GACTCGTCTT ACCTTTCGGG AGCGGACCTC AACCTCTCCT GCCACTCGGC CTCTAACCCA  
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 ATCGCCAAAA TCACGCCAAA TAATAACGGG ACCTATGCCT GTTTTGTCTC TAACTTGGCT  
 ACTGGCCGCA ATAATTCCAT AGTCAAGAGC ATCACAGTCT CTGCATCTGG AACTTCTCCT  
 GGTCTCTCAG CTGGGGCCAC TGTGCGCATC ATGATTGGAG TGCTGGTTGG GGTGCTCTG ATATAG  
 45